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[\[PDF\] Clustering navigation patterns on a website using a sequence alignment method - all 12 versions »](#)

B Hay, G Weits, K Vanhoof - Proc. Intelligent Techniques for Web Personalization: 17th ... 2000 - uhasselt.be

... The Sequence Alignment Method is a non-Euclidean ... unique elements and the costs for reordering common elements. ... are compared with regard to pages and other ...

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[Image management in sport organisations: the creation of value - all 3 versions »](#)

A Ferrand, M Pages - European Journal of Marketing, 1999 - emeraldinsight.com

... Alain Ferrand and Monique Pages Université Claude Bernard ... Juventus should obviously align itself with a company ... choose to reinforce the common elements of the ...

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[A position-sensitive sequence-alignment method illustrated for space-time activity-diary data - all 5 versions »](#)

CH Joh, TA Arentz, HJP Timmermans - Environment and Planning A, 2001 - envplan.com

... deleted) and, as a result, preserved during the alignment. ... if the source sequence has four common elements and the target has two of the same element, then two ...

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[Segmentation of visiting patterns on web sites using a sequence alignment method - all 9 versions »](#)

B Hay, G Weits, K Vanhoof - Journal of Retailing and Consumer Services, 2003 - Elsevier

... information (through the order of elements) within sequences. ... we will introduce the Sequence Alignment Method (SAM ... method is compared to a common used distance ...

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[Sequence alignment with tandem duplication - all 8 versions »](#)

G Benson - Proceedings of the first annual international conference on ..., 1997 - portal.acm.org

... The SI model reflects common mutational events and within this framework, dynamic programming ... scope for two se- quences and to produce an alignment of their ...

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[Querying multiple bioinformatics information sources: can semantic web research help? - all 17 versions »](#)

D Butlier, M Coleman, T Critchlow, R Filato, W Han ... - ACM SIGMOD Record, 2002 - portal.acm.org

... of the DNA sequence around the alignment is extracted to capture the promoter regulatory elements -- the region of ... 15] to find the common transcription binding ...

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[An Empirical Study of the Alignment Between Manufacturing and Marketing Strategies - all 3 versions »](#)

KA Weir, AK Kochhar, SA LeBeau, DG Edgeley - Long Range Planning, 2000 - Elsevier

... a significant part, the infrastructural elements of manufacturing ... use of cross-functional teams, common goals and ... to help cross-functional alignment within ...

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[\[PS\] PUSCA: An Algorithm for Constrained Alignment of Genome Data Sequences - all 10 versions »](#)

I. Marica, A Floratos, I Rigoutsos - Proceedings 9th Workshop on Genome Informatics, 1998 - cs1.cs.nyu.edu

... Similar definitions extend to discover- ing common motifs from multiple sequences. ... 3. Render the alignment. ... Y of subsets of X, such that every element of X ...

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[\[PDF\] Transcription Element Search Software on the WWW - all 4 versions »](#)

J Schug, GC Overton - Technical Report CBIL-TR-1997 - 1001-V 00, of the ..., 1997 - informatik.uni-leipzig.de

... TESS - Transcription Element Search Software, is a software tool ... matrix happens to prefer the less common bases ... The maximumscore possible for any alignment to a ...

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[Managing pages in shared virtual memory systems: getting the compiler into the game](#)

EQ Granston, HAG Wijshoff - Proceedings of the 7th international conference on ..., 1993 - portal.acm.org

... at best. Although program execution is not necessarily interrupted by this, the pages being processed are un- ... A common source of false sharing arises from map ...

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